

Sequence Listing.ST25.txt  
SEQUENCE LISTING

<110> Pfizer Limited  
Bazin, Richard John  
Macdonald, Graeme Arthur  
Phillips, Christopher

<120> Crystal Structure

<130> PCS10934ABXP

<160> 6

<170> PatentIn version 3.0

<210> 1  
<211> 2241  
<212> DNA  
<213> Oryctolagus cuniculus (Rabbit)

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tttggagacgc tggccacctc cccagaaggc acgaggaaaa agcgtttcca aggacggaag 180  
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caccggcgtc tgaagttcct ctcctccaag ttccaggtgc accagatgct caacgagatg 780  
gacgagctga aggagctgaa gaacaaccct caccgcgatt tttacaactg caggaaggtg 840  
gacacccaca tccatgcagc tgcctgcatt aaccagaaac atctgctgcg cttcattaaag 900  
960

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 <211> 747  
 <212> PRT  
 <213> Oryctolagus cuniculus (Rabbit)

<400> 2

Met Pro Leu Phe Lys Leu Pro Ala Glu Gly Lys Glu Leu Asp Asp Ala

## Sequence Listing.ST25.txt

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35 40 45Ile Ser His His Glu Met Gln Ala His Ile Leu His Met Glu Thr Leu  
50 55 60Ala Thr Ser Pro Glu Gly Thr Arg Lys Lys Arg Phe Gln Gly Arg Lys  
65 70 75 80Thr Val Asn Leu Ser Ile Pro Leu Ser Glu Ala Ser Ser Thr Lys Leu  
85 90 95Ser His Ile Asp Glu Tyr Ile Ser Leu Ser Pro Thr Tyr Gln Thr Val  
100 105 110Pro Asp Phe Gln Arg Val Gln Ile Thr Gly Asp Tyr Ala Ser Gly Val  
115 120 125Thr Val Glu Asp Phe Glu Ile Val Cys Lys Gly Leu Tyr Arg Ala Leu  
130 135 140Cys Ile Arg Glu Lys Tyr Met Leu Lys Ser Phe Gln Arg Phe Pro Lys  
145 150 155 160Thr Pro Ser Lys Tyr Leu Arg Ser Ile Glu Gly Thr Ala Trp Lys Ala  
165 170 175Asn Glu Ser Ser Tyr Pro Val Phe Thr Pro Ala Leu Lys Lys Gly Glu  
180 185 190Asp Pro Phe Arg Thr Asp Asn Leu Pro Glu Asn Leu Gly Tyr His Leu  
195 200 205Lys Met Lys Asp Gly Val Val Tyr Ile Tyr Ala Asn Glu Ala Ala Ala  
210 215 220Gly Lys Asp Glu Pro Lys Pro Leu Leu Tyr Pro Asn Met Glu Glu Phe  
225 230 235 240Leu Asp Asp Met Asn Phe Leu Leu Ala Leu Ile Ala Gln Gly Pro Val  
245 250 255Lys Thr Tyr Thr His Arg Arg Leu Lys Phe Leu Ser Ser Lys Phe Gln  
260 265 270Val His Gln Met Leu Asn Glu Met Asp Glu Leu Lys Glu Leu Lys Asn  
275 280 285

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Asn Pro His Arg Asp Phe Tyr Asn Cys Arg Lys Val Asp Thr His Ile  
290 295 300

His Ala Ala Ala Cys Met Asn Gln Lys His Leu Leu Arg Phe Ile Lys  
305 310 315 320

Lys Ser Tyr Gln Val Asp Ala Asp Arg Val Val Tyr Ser Thr Lys Glu  
325 330 335

Lys Asn Leu Thr Leu Lys Gln Leu Phe Asp Lys Leu Lys Leu His Pro  
340 345 350

Tyr Asp Leu Thr Val Asp Ser Leu Asp Val His Ala Gly Arg Gln Thr  
355 360 365

Phe Gln Arg Phe Asp Lys Phe Asn Asp Lys Tyr Asn Pro Val Gly Ala  
370 375 380

Ser Glu Leu Arg Asp Leu Tyr Leu Lys Thr Asp Asn Tyr Ile Asn Gly  
385 390 395 400

Glu Tyr Phe Ala Thr Ile Ile Lys Glu Val Gly Ala Asp Leu Val Asp  
405 410 415

Ala Lys Tyr Gln His Ala Glu Pro Arg Leu Ser Ile Tyr Gly Arg Ser  
420 425 430

Pro Asp Glu Trp Ser Lys Leu Ser Ser Trp Phe Val Arg Asn Arg Ile  
435 440 445

Tyr Ser Ser Asn Met Thr Trp Met Ile Gln Val Pro Arg Ile Tyr Asp  
450 455 460

Val Phe Arg Ser Lys Asn Phe Leu Pro His Phe Gly Lys Met Leu Glu  
465 470 475 480

Asn Val Phe Met Pro Val Phe Glu Ala Thr Ile Asn Pro Gln Ala His  
485 490 495

Pro Glu Leu Ser Val Phe Leu Lys His Ile Thr Gly Phe Asp Ser Val  
500 505 510

Asp Asp Glu Ser Lys His Ser Gly His Met Phe Ser Ser Lys Ser Pro  
515 520 525

Lys Pro Gln Glu Trp Thr Leu Glu Lys Asn Pro Ser Tyr Thr Tyr Tyr  
530 535 540

Ala Tyr Tyr Met Tyr Ala Asn Ile Met Val Leu Asn Ser Leu Arg Lys  
545 550 555 560

Glu Arg Gly Met Asn Thr Phe Leu Phe Arg Pro His Cys Gly Glu Val  
565 570 575

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Gly Ala Leu Thr His Leu Met Thr Ala Phe Met Thr Ala Asp Asn Ile  
580 585 590

Ser His Gly Leu Asn Leu Lys Lys Ser Pro Val Leu Gln Tyr Leu Phe  
595 600 605

Phe Leu Ala Gln Ile Pro Ile Ala Met Ser Pro Leu Ser Asn Asn Ser  
610 615 620

Leu Phe Leu Glu Tyr Ala Lys Asn Pro Phe Leu Asp Phe Leu Gln Lys  
625 630 635 640

Gly Leu Met Ile Ser Leu Ser Thr Asp Asp Pro Met Gln Phe His Phe  
645 650 655

Thr Lys Glu Pro Leu Met Glu Glu Tyr Ala Ile Ala Ala Gln Val Phe  
660 665 670

Lys Leu Ser Thr Cys Asp Met Cys Glu Val Ala Arg Asn Ser Val Leu  
675 680 685

Gln Cys Gly Ile Ser His Glu Glu Lys Ala Lys Phe Leu Gly Asn Asn  
690 695 700

Tyr Leu Glu Glu Gly Pro Ile Gly Asn Asp Ile Arg Lys Thr Asn Val  
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Ala Gln Ile Arg Met Ala Tyr Arg Tyr Glu Thr Trp Cys Tyr Glu Leu  
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Asn Leu Ile Ala Glu Gly Leu Lys Ser Thr Glu  
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<210> 3

<211> 20

<212> DNA

<213> Homo sapiens

<400> 3

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20

<210> 4

<211> 21

<212> DNA

<213> Homo sapiens

<400> 4

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21

<210> 5

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<211> 21  
<212> DNA  
<213> Homo sapiens

<400> 5  
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<210> 6  
<211> 22  
<212> DNA  
<213> Homo sapiens

<400> 6  
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